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CLAIMS IN THE CASE

- (currently amended) A process of preparing unsaturated fatty acids, which
 comprises introducing, into an organism, at least one isolated nucleic acid
 sequence encoding a polypeptide having 6-desaturase activity, selected from
 the group consisting of:
 - a) a nucleic acid sequence having the sequence shown in SEQ ID NO: 1,
 - b) nucleic acid sequences which, as a result of the degeneracy of the genetic code, are derived from the [lacuna] sequence shown in SEQ ID NO: 1,
 - c) derivatives of the nucleic acid sequence shown in SEQ ID NO: 1 which encode polypeptides with the amino acid sequences shown in SEQ ID NO: 2 and have at least 50% homology at the amino acid level without substantially reducing the enzymatic action of the polypeptides,

and culturing this organism, where the cultured organism contains at least 1 mol% of unsaturated fatty acids based on the total fatty acid content in the organism.

- (original) The process as claimed in claim 1, wherein the nucleic acid sequence is derived from a plant or algae.
- (previously presented) The process as claimed in claim 1, wherein the nucleic acid sequence is derived from Physcomitrella patens.
- (previously presented) The process as claimed in claim 1, wherein the organism
 is an organism selected from the group consisting of bacterium, fungus, ciliate,

HEINZ et al., Ser. No. 10,019,048 algae, cyanobacterium, animal or plant.

- (previously presented) The process as claimed in claim 1, wherein the organism is a plant or algae.
- 6. (previously presented) The process as claimed in claim 1, wherein the organism is an oil crop.
- 7. (previously presented) The process as claimed in claim 1, wherein the cultured organism contains at least 5% by weight of unsaturated fatty acids based on the total fatty acid content in the organism.
- (previously presented) The process as claimed in claim 1, wherein the unsaturated fatty acids are isolated from the organism.
- 9. (original) A transgenic organism selected from the group consisting of plants, fungi, ciliates, algae, bacteria, cyanobacteria or animals comprising at least one isolated nucleic acid sequence encoding a polypeptide with 6-desaturase activity, selected from the group consisting of:
 - a) a nucleic acid sequence having the sequence shown in SEQ ID NO: 1,
 - b) nucleic acid sequences which, as a result of the degeneracy of the genetic code, are derived from the sequence shown in SEQ ID NO: 1,
 - c) derivatives of the nucleic acid sequence shown in SEQ ID NO: 1 which encode polypeptides with the amino acid sequences shown in SEQ ID NO: 2 and have at least 50% homology at the amino acid level without substantially reducing the enzymatic action of the polypeptides.
- 10. (original) A transgenic organism as claimed in claim 9, wherein the organism is a

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- 11. (withdrawn)
- 12. (withdrawn)